

# PCT 371 Routing Sheet

## APPLICATION

10/554,240

IFW DocCode- SEQREQ

Index using Current Date

### TO BE DELIVERED TO:

Tech Center Scanning

### Sequence Rule Compliance Review Item

	CRF, paper copy of sequence listing, and statement that both are same missing
	CRF contains error(s) according to STIC Report
	CRF damaged or unreadable according to STIC Report
	CRF transferred from prior application is not compliant
X	OTHER

Place an "X" in the appropriate box



Cheryl Tseng  
Patent Examiner  
Technology Center 1000

Comment Sheet

**APPLICATION SERIAL NUMBER**

**10/554,240**

**DOES NOT COMPLY WITH THE  
SEQUENCE RULES. See reasons below.**

The claims and specification recite sequences which are embraced by the sequence rules. Applicant's submission filed November 27, 2006 did not fully comply with the requirement. For example, the sequences found in ¶[0018], lines 2-9 recite sequences which require sequence identifiers. Compliance with 37 CFR 1.821-1.825 excludes only branched sequences, sequences with fewer than 4 specifically defined amino acids and sequences with D-amino acids, however these exemplary sequences do not fall within the exclusions. Further, the sequences found in claim 19 require Sequence identifiers for the same reasons. 37 CFR 1.821(a)(2) states (in part), Any peptide or protein that can be expressed as a sequence using the symbols in WIPO Standard ST.25 (1998), Appendix 2, Table 3 in conjunction with a description in the Feature section to describe, for example, modified linkages, cross links and end caps, non-peptidyl bonds, etc., is embraced by this definition.